

Additional file 2: Supplementary tables S1 and S2

Development of a novel heterologous β -lactam-specific whole-cell biosensor in *Bacillus subtilis*

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Table S2: Vector backbones and expression vectors used and designed in this study.

Table S1: Primers used in this study.

Primer #	Primer name	Primer sequence 5'-3'	Reference
Primers to amplify genetic regions of interest			
TM5006	<i>blaR1</i> -check-1	CTGTACCTGTTTTCCCATACAGT	this study
TM5007	<i>blaR1</i> -check-2	GGATTGCCCCATTAAAACTGG	this study
TM5008	<i>blaI</i> -rev	GATCGAATTCGCGGCCGCTTCTAGATTACTTTTTACTA ATATC	this study
TM5009	<i>blaR1</i> -fwd-sall	GATCGTCGACATATTACAGTTGTAATTTTT	this study
TM5010	<i>blaR1</i> -fwd-suffix	GATCCTGCAGCGGCCGCTACTAGTATATTACAGTTGTA ATTTTT	this study
iG17P035	P _{blaZ} fwd RFC10 (syn)	<u>GATCGAATTCGCGGCCGCTTCTAGATTCAAATATTTAT</u> AATAACAATTGACATCAATATTACAATTGTAATATTATT G	this study
iG17P036	P _{blaZ} rev RFC 10 (syn)	<u>GATCCTGCAGCGGCCGCTACTAGTATATTACAGTTGTA</u> ATTTTTATAAATCAATAATATTACAATTGTAATATTGATG	this study
iG17P037	P _{blaR1} fwd RFC10 (syn)	<u>GATCGAATTCGCGGCCGCTTCTAGAATATTACAGTTGT</u> AATTTTTATAAATCAATAATATTACAATTGTAATATTGAT G	this study
iG17P038	P _{blaR1} rev RFC10 (syn)	<u>GATCCTGCAGCGGCCGCTACTAGTTTCAAATATTTATA</u> ATAACAATTGACATCAATATTACAATTGTAATATTATT G	this study
iG17P039	Biobrick_suffix_rev	GATCCTGCAGCGGCCGCTACTAGT	this study
iG17P040	P _{penP} fwd up RFC10 long	<u>GATCGGAATTCGCGGCCGCTTCTAGATCTGGTTTTATG</u> CTTAATCCTC	this study
iG17P041	P _{penP} rev RFC10	<u>GATCCTGCAGCGGCCGCTACTAGTTCAAATGATTTTGT</u> ATTACCTTTG	this study
iG17P042	P _{penP} fwd up RFC10 short	<u>GATCGGAATTCGCGGCCGCTTCTAGAAATCACAATTGA</u> TAAAGCTTTCTAA	this study
iG17P043	<i>penP</i> fwd RFC25	<u>GATCGAATTCGCGGCCGCTTCTAGATCAGAGGAGGC</u> <u>CTGATGGCCGGCAAGTTGAAAATAAAGCGTCAATAA</u> <u>GATCCTGCAGCGGCCGCTACTAGTATTAACCGGTTTT</u>	this study
iG17P044	<i>penP</i> rev RFC25	<u>GAGATCGTTAAGGACGAC</u> <u>GATCGAATTCGCGGCCGCTTCTAGATAGGAGGTGTC</u> <u>AAAATGGCCGGCGCCAACTGCTCATTATGTC</u>	this study
iG17P153	<i>blaR1</i> fwd RFC25	<u>GATCCTGCAGCGGCCGCTACTAGTATTAACCGGTTTTG</u> GTCGTTCAAAACACCC	this study
iG17P154	<i>blaR1</i> rev RFC25	<u>GATCCTGCAGCGGCCGCTACTAGTTTTGAGATCGTTAA</u> GGACGAC	this study
iG17P227	<i>penP</i> rev RFC10 w/o stop	<u>GATCCTGCAGCGGCCGCTACTAGTTTTGAGATCGTTAA</u> GGACGAC	this study
iG17P228	<i>blaR1</i> rev RFC10 w/o stop	<u>GATCCTGCAGCGGCCGCTACTAGTTTTGGTCGTTCAAA</u> ACACCCATTC	this study
TM4487	Biobrick_prefix_fwd	GATCGAATTCGCGGCCGCTTCTAGA	
TM5136	<i>penP</i> -up_fwd	GGGTCTTTTCTGATCAAGACAGTC	this study
TM5137	<i>penP</i> -up_rev	CCTATCACCTCAAATGGTTCGCTGGCCCAACACATATT CCGAATTTTATTG	this study
TM5138	<i>penP</i> -do_fwd	CGAGCGCCTACGAGGAATTTGTATCG CTGAGGCTGCAAAAGTCGTCC	this study
TM5139	<i>penP</i> -do_rev	GCTGGAATCCAGAAAGAAGCGG	this study
TM5140	<i>ybxI</i> -up_fwd	GCATGCTGTATACACTGGGTGTC	this study
TM5141	<i>ybxI</i> -up_rev	CCTATCACCTCAAATGGTTCGCTGCACAAAAGCCAAAA ACATACCGTG	this study
TM5142	<i>ybxI</i> -do_fwd	CGAGCGCCTACGAGGAATTTGTATCGCCTGCAATACT CAGCACAAGCAC	this study
TM5143	<i>ybxI</i> -do_rev	GACCTCCTGCACAAACATTTCTC	this study

Primers used to check for integration into vector backbone		
TM2262	pAH328checkfwd	GAGCGTAGCGAAAAATCC
TM2263	pAH328checkrev	GAAATGATGCTCCAGTAACC
TM3081	pSBBs2E seq fwd	GGCAACCGAGCGTTCTG
TM3082	pSBBs2E seq rev	CTGACAGCGTTTCGATCC
TM0747	thrC-check-fwd	CGCTCAAGCTGTCATGTACG
TM0149	spec-check rev	CGTATGTATTCAAATATATCCTCCTCAC
Primers to prove integration into <i>Bacillus subtilis</i> genome		
TM4085	pBS2E int. up fwd	TGCTGCAAAAGAATTTTGTGTCCG
TM4086	pBS2E int. up rev	AGGACTCTCTAGCTTGAGGC
TM4087	pBS2E int. do fwd	CTGCAGAGATATCGATTTC AAGC
TM4088	pBS2E int. do rev	CTTTGCTTTTCATGATTTCATCCC
TM2505	pAH328 sacA front check fwd	CTGATTGGCATGGCGATTGC
TM2506	pAH328 sacA front check rev	ACAGCTCCAGATCCTCTACG
TM2507	pAH328 sacA back check fwd	GTCGCTACCATTACCA GTTG
TM2508	pAH328 sacA back check rev	TCCAAACATTCCGGTGTTATC

The Primer number (#) and name refer to the number and the name specified in the primer list of the research group. The primer sequences are displayed in 5' to 3' direction. Underlined sequences indicate overhangs with endonuclease recognition sites.

Table S2: Vector backbones and expression vectors used and designed in this study.

Vector	Description	Source	Reference
Vector backbones used in all experiments			
pBS2E	empty vector, integration into <i>lacA</i> , <i>amp^r</i> , <i>mls^r</i>	pAX01-derivative	Radeck <i>et al.</i> 2013
pBS4S	empty vector, integration into <i>thrC</i> , <i>amp^r</i> , <i>spec^r</i>	pDG1731-derivative	Radeck <i>et al.</i> 2013
pBS3C<i>lux</i>	<i>lux</i> -reporter vector, integration into <i>sacA</i> , <i>amp^r</i> , <i>cm^r</i>	pAH328-derivative	Radeck <i>et al.</i> 2013
Vectors developed in this study			
pBS3C-P_{blaR1}-<i>blaR1</i>-P_{blaZ}-<i>lux</i> (ExSall)	<i>lux</i> -reporter vector, integration into <i>sacA</i> , <i>amp^r</i> , <i>cm^r</i>	pAH328-derivative	This study
pBS2E-P_{xyIA}-<i>blaR1</i>	P _{xyIA} - <i>blaR1</i> in MCS, integration into <i>lacA</i> , <i>amp^r</i> , <i>mls^r</i>	pAX01-derivative	This study
pBS2E-P_{veg}-<i>blaR1</i>	P _{veg} - <i>blaR1</i> in MCS, integration into <i>lacA</i> , <i>amp^r</i> , <i>mls^r</i>	pAX01-derivative	This study
pBS4S-P_{lepA}-<i>blaI</i>	P _{lepA} - <i>blaI</i> in MCS, integration into <i>thrC</i> , <i>amp^r</i> , <i>spec^r</i>	pDG1731-derivative	This study
pBS3C-P_{blaZ}-<i>lux</i>	P _{blaZ} in MCS, <i>lux</i> -reporter vector, integration into <i>sacA</i> , <i>amp^r</i> , <i>cm^r</i>	pAH328-derivative	This study